



22	19	100.0	422	18	95	09	132	244	2733	Suspense	2733	Ap	17	89.5	602	31	03	09	645	294	47698	Suspense	47698	Ap	17	89.5	602	31	03	09	645	294	47698
23	19	100.0	439	25	05	09	652	293	4769	Suspense	4769	Ap	17	89.5	7141	1	03	09	645	294	47698	Suspense	47698	Ap	17	89.5	7141	1	03	09	645	294	47698
24	19	100.0	441	25	05	09	652	122	138	Suspense	138	Ap	17	89.5	7141	1	03	09	645	294	47698	Suspense	47698	Ap	17	89.5	7141	1	03	09	645	294	47698
25	19	100.0	443	25	05	09	652	455	4711	Suspense	4711	Ap	17	89.5	7141	1	03	09	645	294	47698	Suspense	47698	Ap	17	89.5	7141	1	03	09	645	294	47698
26	19	100.0	449	17	08	09	439	740	2779	Suspense	2779	Ap	17	89.5	7141	1	03	09	645	294	47698	Suspense	47698	Ap	17	89.5	7141	1	03	09	645	294	47698
27	19	100.0	449	44	08	09	921	478	2179	Suspense	2179	Ap	17	89.5	7141	1	03	09	645	294	47698	Suspense	47698	Ap	17	89.5	7141	1	03	09	645	294	47698
28	19	100.0	459	17	08	09	439	087	3008	Suspense	3008	Ap	17	89.5	7141	1	03	09	645	294	47698	Suspense	47698	Ap	17	89.5	7141	1	03	09	645	294	47698
29	19	100.0	459	25	05	09	721	588	524	Suspense	524	Ap	102	102	645	294	47698	Suspense	47698	Ap	102	102	645	294	47698	Suspense	47698	Ap	102	102	645	294	47698
30	19	100.0	452	27	03	09	659	530	2174	Suspense	2179	Ap	163	163	47698	1	03	09	645	294	47698	Suspense	47698	Ap	163	163	47698	1	03	09	645	294	47698
31	19	100.0	485	27	03	09	659	908	2174	Suspense	2179	Ap	163	163	47698	1	03	09	645	294	47698	Suspense	47698	Ap	163	163	47698	1	03	09	645	294	47698
32	19	100.0	494	17	03	09	473	196	2433	Suspense	2433	Ap	163	163	47698	1	03	09	645	294	47698	Suspense	47698	Ap	163	163	47698	1	03	09	645	294	47698
33	19	100.0	494	41	03	09	803	853	2433	Suspense	2433	Ap	163	163	47698	1	03	09	645	294	47698	Suspense	47698	Ap	163	163	47698	1	03	09	645	294	47698
34	19	100.0	504	29	03	09	726	175	1968	Suspense	1968	Ap	163	163	47698	1	03	09	645	294	47698	Suspense	47698	Ap	163	163	47698	1	03	09	645	294	47698
35	19	100.0	506	29	03	09	726	455	524	Suspense	524	Ap	163	163	47698	1	03	09	645	294	47698	Suspense	47698	Ap	163	163	47698	1	03	09	645	294	47698
36	19	100.0	506	42	03	10	222	609	524	Suspense	524	Ap	163	163	47698	1	03	09	645	294	47698	Suspense	47698	Ap	163	163	47698	1	03	09	645	294	47698
37	19	100.0	540	18	05	09	432	241A	2511	Suspense	2084	Ap	113	113	645	294	47698	Suspense	47698	Ap	113	113	645	294	47698	Suspense	47698	Ap	113	113	645	294	47698
38	19	100.0	552	25	05	09	649	162	2084	Suspense	2084	Ap	113	113	645	294	47698	Suspense	47698	Ap	113	113	645	294	47698	Suspense	47698	Ap	113	113	645	294	47698
39	19	100.0	603	42	05	10	406	021	131	Suspense	171	Ap	112	112	645	294	47698	Suspense	47698	Ap	112	112	645	294	47698	Suspense	47698	Ap	112	112	645	294	47698
40	19	100.0	654	29	03	09	721	589	496	Suspense	496	Ap	113	113	645	294	47698	Suspense	47698	Ap	113	113	645	294	47698	Suspense	47698	Ap	113	113	645	294	47698
41	19	100.0	763	67	02	06	244	690	10610	Suspense	10610	A	114	114	645	294	47698	Suspense	47698	Ap	114	114	645	294	47698	Suspense	47698	Ap	114	114	645	294	47698
42	19	100.0	763	76	02	06	244	690	10610	Suspense	10610	A	114	114	645	294	47698	Suspense	47698	Ap	114	114	645	294	47698	Suspense	47698	Ap	114	114	645	294	47698
43	19	100.0	774	45	03	10	213	843	222	Suspense	222	Ap	115	115	645	294	47698	Suspense	47698	Ap	115	115	645	294	47698	Suspense	47698	Ap	115	115	645	294	47698
44	19	100.0	1062	17	03	09	496	087	3008	Suspense	3008	Ap	116	116	645	294	47698	Suspense	47698	Ap	116	116	645	294	47698	Suspense	47698	Ap	116	116	645	294	47698
45	19	100.0	1265	25	05	09	652	122	4769	Suspense	4769	Ap	118	118	645	294	47698	Suspense	47698	Ap	118	118	645	294	47698	Suspense	47698	Ap	118	118	645	294	47698
46	19	100.0	1265	25	05	09	726	733	524	Suspense	524	Ap	119	119	645	294	47698	Suspense	47698	Ap	119	119	645	294	47698	Suspense	47698	Ap	119	119	645	294	47698
47	19	100.0	1745	28	05	09	705	256A	7619	Suspense	7619	Ap	120	120	645	294	47698	Suspense	47698	Ap	120	120	645	294	47698	Suspense	47698	Ap	120	120	645	294	47698
48	19	100.0	1745	60	05	10	164	285	7619	Suspense	7619	Ap	121	121	645	294	47698	Suspense	47698	Ap	121	121	645	294	47698	Suspense	47698	Ap	121	121	645	294	47698
49	19	100.0	1760	45	03	09	646	076A	131	Suspense	131	Ap	122	122	645	294	47698	Suspense	47698	Ap	122	122	645	294	47698	Suspense	47698	Ap	122	122	645	294	47698
50	19	100.0	1760	43	03	10	131	430	131	Suspense	131	Ap	123	123	645	294	47698	Suspense	47698	Ap	123	123	645	294	47698	Suspense	47698	Ap	123	123	645	294	47698
51	19	100.0	1761	71	03	09	478	258	4195	Suspense	4195	Ap	124	124	645	294	47698	Suspense	47698	Ap	124	124	645	294	47698	Suspense	47698	Ap	124	124	645	294	47698
52	19	100.0	1764	17	03	09	359	522	455	Suspense	455	Ap	125	125	645	294	47698	Suspense	47698	Ap	125	125	645	294	47698	Suspense	47698	Ap	125	125	645	294	47698
53	19	100.0	1764	17	03	09	359	522	455	Suspense	455	Ap	126	126	645	294	47698	Suspense	47698	Ap	126	126	645	294	47698	Suspense	47698	Ap	126	126	645	294	47698
54	19	100.0	1764	41	03	09	919	092	455	Suspense	455	Ap	127	127	645	294	47698	Suspense	47698	Ap	127	127	645	294	47698	Suspense	47698	Ap	127	127	645	294	47698
55	19	100.0	1786	11	03	08	752	345	1	Suspense	1	Ap	128	128	645	294	47698	Suspense	47698	Ap	128	128	645	294	47698	Suspense	47698	Ap	128	128	645	294	47698
56	19	100.0	1786	14	03	09	015	434	131	Suspense	131	Ap	129	129	645	294	47698	Suspense	47698	Ap	129	129	645	294	47698	Suspense	47698	Ap	129	129	645	294	47698
57	19	100.0	1789	1	03	10	050A	4136A	131	Suspense	131	Ap	130	130	645	294	47698	Suspense	47698	Ap	130	130	645	294	47698	Suspense	47698	Ap	130	130	645	294	47698
58	19	100.0	1789	14	03	09	053	473A	131	Suspense	131	Ap	131	131	645	294	47698	Suspense	47698	Ap	131	131	645	294	47698	Suspense	47698	Ap	131	131	645	294	47698
59	19	100.0	1789	14	03	09	053	473A	131	Suspense	131	Ap	132	132	645	294	47698	Suspense	47698	Ap	132	132	645	294	47698	Suspense	47698	Ap	132	132	645	294	47698
60	19	100.0	1789	22	05	09	589	034	1	Suspense	1	Ap	133	133	645	294	47698	Suspense	47698	Ap	133	133	645	294	47698	Suspense	47698	Ap	133	133	645	294	47698
61	19	100.0	1789	32	05	09	840	704	1	Suspense	1	Ap	134	134	645	294	47698	Suspense	47698	Ap	134	134	645	294	47698	Suspense	47698	Ap	134	134	645	294	47698
62	19	100.0	1789	36	03	09	967	854	1	Suspense	1	Ap	135	135	645	294	47698	Suspense	47698	Ap	135	135	645	294	47698	Suspense	47698	Ap	135	135	645	294	47698
63	19	100.0	1789	45	03	10	111	534	1	Suspense	1	Ap	136	136	645	294	47698	Suspense	47698	Ap	136	136	645	294	47698	Suspense	47698	Ap	136	136	645	294	47698
64	19	100.0	1789	41	03	10	188	893	1	Suspense	1	Ap	137	137	645	294	47698	Suspense	47698	Ap	137	137	645	294	47698	Suspense	47698	Ap	137	137	645	294	47698
65	19	100.0	1801	40	03	09	760	444	691	Suspense	691	Ap	138	138	645	294	47698	Suspense	47698	Ap	138	138	645	294	47698	Suspense	47698	Ap	138	138	645	294	47698
66	19	100.0	1801	42	03	10	206	021	241	Suspense	294	Ap	139	139	645	294	47698	Suspense	47698	Ap	139	139	645	294	47698	Suspense	47698	Ap	139	139	645	294	47698
67	19	100.0	1801	42	03	10	212	074	491	Suspense																							

16.4	86.3	17427	58	US-60-144	551-549	Sequence 549, App	241	84.2	1515	27	US-09-684	016-385492	16.4	86.3	17427	58	US-60-144	551-549	Sequence 549, App	241	84.2	1515	27	US-09-684	016-385492
16.4	86.3	17995	24	US-09-620-692-17124		Sequence 17124, A	242	84.2	1524	41	US-10-155	881-35405	16.4	86.3	17995	24	US-09-620-692-17124		Sequence 17124, A	242	84.2	1524	41	US-10-155	881-35405
16.4	86.3	18540	24	US-09-620-692-4542		Sequence 448, App	243	84.2	1524	75	US-60-812-544-524		16.4	86.3	18540	24	US-09-620-692-4542		Sequence 448, App	243	84.2	1524	75	US-60-812-544-524	
16.4	86.3	18540	28	US-09-702-134-29462		Sequence 29462, A	244	84.2	2112	41	US-10-155	881-18543	16.4	86.3	18540	28	US-09-702-134-29462		Sequence 29462, A	244	84.2	2112	41	US-10-155	881-18543
16.4	86.3	18540	31	US-09-615-264-74324		Sequence 74324, A	245	84.2	22916	1	US-10-155	881-20117	16.4	86.3	18540	31	US-09-615-264-74324		Sequence 74324, A	245	84.2	22916	1	US-10-155	881-20117
16.4	86.3	18611	24	US-09-620-392-45167		Sequence 45167, A	246	84.2	22916	40	US-09-764-906-20117		16.4	86.3	18611	24	US-09-620-392-45167		Sequence 45167, A	246	84.2	22916	40	US-09-764-906-20117	
16.4	86.3	18611	28	US-09-702-134-63430		Sequence 63430, App	247	84.2	22916	47	US-10-155	881-179264	16.4	86.3	18611	28	US-09-702-134-63430		Sequence 63430, App	247	84.2	22916	47	US-10-155	881-179264
16.4	86.3	18611	31	US-09-615-264-75498		Sequence 75498, A	248	84.2	1243197	24	US-60-212-664-3		16.4	86.3	18611	31	US-09-615-264-75498		Sequence 75498, A	248	84.2	1243197	24	US-60-212-664-3	
16.4	86.3	22119	21	US-09-620-702-64098		Sequence 64098, A	249	84.2	1243197	48	US-10-155	881-179264	16.4	86.3	22119	21	US-09-620-702-64098		Sequence 64098, A	249	84.2	1243197	48	US-10-155	881-179264
16.4	86.3	25144	24	US-09-620-392-64098		Sequence 64098, A	250	84.2	1243197	48	US-10-155	881-179264	16.4	86.3	25144	24	US-09-620-392-64098		Sequence 64098, A	250	84.2	1243197	48	US-10-155	881-179264
16.4	86.3	25144	28	US-09-702-134-51576		Sequence 51576, A	251	84.2	1243197	12	US-10-155	881-179264	16.4	86.3	25144	28	US-09-702-134-51576		Sequence 51576, A	251	84.2	1243197	12	US-10-155	881-179264
16.4	86.3	25144	31	US-09-615-264-75498		Sequence 75498, A	252	84.2	1243197	12	US-10-155	881-179264	16.4	86.3	25144	31	US-09-615-264-75498		Sequence 75498, A	252	84.2	1243197	12	US-10-155	881-179264
16.4	86.3	25144	34	US-09-620-392-63430		Sequence 63430, App	253	84.2	1243197	12	US-10-155	881-179264	16.4	86.3	25144	34	US-09-620-392-63430		Sequence 63430, App	253	84.2	1243197	12	US-10-155	881-179264
16.4	86.3	25144	37	US-09-620-392-63430		Sequence 63430, App	254	84.2	1243197	12	US-10-155	881-179264	16.4	86.3	25144	37	US-09-620-392-63430		Sequence 63430, App	254	84.2	1243197	12	US-10-155	881-179264
16.4	86.3	25144	40	US-09-620-392-63430		Sequence 63430, App	255	84.2	1243197	12	US-10-155	881-179264	16.4	86.3	25144	40	US-09-620-392-63430		Sequence 63430, App	255	84.2	1243197	12	US-10-155	881-179264
16.4	86.3	25197	28	US-09-702-134-15994		Sequence 25904, A	256	84.2	1243197	184	US-60-812-544-524		16.4	86.3	25197	28	US-09-702-134-15994		Sequence 25904, A	256	84.2	1243197	184	US-60-812-544-524	
16.4	86.3	25197	31	US-09-615-264-75498		Sequence 75498, A	257	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	25197	31	US-09-615-264-75498		Sequence 75498, A	257	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	34107	24	US-09-620-392-66588		Sequence 66588, A	258	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	34107	24	US-09-620-392-66588		Sequence 66588, A	258	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	34111	28	US-09-702-134-5064		Sequence 5064, App	259	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	34111	28	US-09-702-134-5064		Sequence 5064, App	259	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	34111	31	US-09-615-264-75498		Sequence 63381, A	260	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	34111	31	US-09-615-264-75498		Sequence 63381, A	260	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	34111	34	US-09-620-392-63430		Sequence 45610, A	261	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	34111	34	US-09-620-392-63430		Sequence 45610, A	261	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	34735	24	US-09-620-392-45610		Sequence 45610, A	262	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	34735	24	US-09-620-392-45610		Sequence 45610, A	262	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	41240	24	US-09-620-392-14275		Sequence 11, App1	263	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	41240	24	US-09-620-392-14275		Sequence 11, App1	263	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	65669	68	US-60-245-224-11		Sequence 6, App1	264	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	65669	68	US-60-245-224-11		Sequence 6, App1	264	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	66379	28	US-09-702-134-6347		Sequence 6347, App	265	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	66379	28	US-09-702-134-6347		Sequence 6347, App	265	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	66379	31	US-09-615-264-75498		Sequence 80437, A	266	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	66379	31	US-09-615-264-75498		Sequence 80437, A	266	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	89740	28	US-09-620-392-4845		Sequence 89740, App	267	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	89740	28	US-09-620-392-4845		Sequence 89740, App	267	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	89740	31	US-09-615-264-75498		Sequence 236, App	268	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	89740	31	US-09-615-264-75498		Sequence 236, App	268	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	14488	65	US-60-212-664-246		Sequence 9705, A	269	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	14488	65	US-60-212-664-246		Sequence 9705, A	269	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	165972	28	US-09-702-134-9705		Sequence 65753, A	270	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	165972	28	US-09-702-134-9705		Sequence 65753, A	270	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	165972	31	US-09-615-264-75498		Sequence 128, App	271	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	165972	31	US-09-615-264-75498		Sequence 128, App	271	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312457	35	US-09-948-947-128		Sequence 157, App	272	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	312457	35	US-09-948-947-128		Sequence 157, App	272	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	273	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	273	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	274	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	274	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	275	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	275	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	276	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	276	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	277	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	277	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	278	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	278	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	279	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	279	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	280	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	280	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	281	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	281	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	282	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	282	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	283	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	283	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	284	84.2	1243197	236	US-09-250-210A-5043		16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	284	84.2	1243197	236	US-09-250-210A-5043	
16.4	86.3	312472	35	US-09-948-947-157		Sequence 532, App	285																		

[illegible]

c 400	15.4	81.1	277	21	US-09-540-229-179781	Sequence 179781, A	534	15.4	81.1	521	15	US-09-297-618-24664	Sequence 24664, A
c 401	15.4	81.1	277	68	US-10-029-666-14031	Sequence 14031, A	c 534	15.4	81.1	521	27	US-09-542-565A-4210	Sequence 4210, A
c 402	15.4	81.1	290	16	US-09-294-094A-464	Sequence 464, App	c 535	15.4	81.1	525	29	US-09-666-455A-5224	Sequence 5224, A
c 403	15.4	81.1	290	16	US-09-294-094B-298	Sequence 298, App	c 536	15.4	81.1	525	29	US-10-029-565-5291	Sequence 5291, A
c 404	15.4	81.1	290	32	US-09-839-976-298	Sequence 298, App	c 537	15.4	81.1	527	29	US-09-668-632-1141	Sequence 1141, A
c 405	15.4	81.1	290	42	US-09-839-976A-298	Sequence 298A, App	c 538	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 406	15.4	81.1	290	52	US-60-082-567-298	Sequence 298, App	c 539	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 407	15.4	81.1	291	25	US-09-649-163-5482	Sequence 5482, App	c 540	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 408	15.4	81.1	291	25	US-09-652-911-5275	Sequence 5275, App	c 541	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 409	15.4	81.1	291	25	US-09-652-911-5275	Sequence 5275, App	c 542	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 410	15.4	81.1	298	17	US-09-865-419A-10140	Sequence 10140, A	c 543	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 411	15.4	81.1	298	17	US-09-865-419B-14666	Sequence 14666, A	c 544	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 412	15.4	81.1	319	17	US-09-432-782-7029	Sequence 7029, App	c 545	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 413	15.4	81.1	319	19	US-09-515-694-7029	Sequence 7029, App	c 546	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 414	15.4	81.1	320	27	US-09-692-257A-3311	Sequence 3311, App	c 547	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 415	15.4	81.1	320	60	US-60-162-747-3451	Sequence 3451, App	c 548	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 416	15.4	81.1	326	17	US-09-442-782-7030	Sequence 7030, App	c 549	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 417	15.4	81.1	326	19	US-09-515-694-7040	Sequence 7040, App	c 550	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 418	15.4	81.1	329	30	US-09-770-791-797	Sequence 797, App	c 551	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 419	15.4	81.1	329	30	US-09-770-791-797	Sequence 797, App	c 552	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 420	15.4	81.1	339	33	US-09-865-439A-7008	Sequence 7008, App	c 553	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 421	15.4	81.1	339	33	US-09-865-439B-106970	Sequence 106970, App	c 554	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 422	15.4	81.1	340	27	US-09-684-616-106970	Sequence 106970, App	c 555	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 423	15.4	81.1	342	17	US-09-304-745-39030	Sequence 39030, A	c 556	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 424	15.4	81.1	342	22	US-09-565-306-57862	Sequence 57862, A	c 557	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 425	15.4	81.1	350	19	US-09-332-782-3506	Sequence 3506, App	c 558	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 426	15.4	81.1	350	19	US-09-515-694-3506	Sequence 3506, App	c 559	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 427	15.4	81.1	356	17	US-09-359-067-32494	Sequence 32494, A	c 560	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 428	15.4	81.1	357	17	US-09-362-510-17212	Sequence 17212, A	c 561	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 429	15.4	81.1	357	17	US-09-362-510A-17212	Sequence 17212, A	c 562	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 430	15.4	81.1	357	34	US-09-904-013-17212	Sequence 12409, A	c 563	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 431	15.4	81.1	377	16	US-09-287-618-12409	Sequence 12409, A	c 564	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 432	15.4	81.1	378	24	US-09-637-623-111	Sequence 111, App	c 565	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 433	15.4	81.1	385	25	US-09-606-776-2051	Sequence 2051, App	c 566	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 434	15.4	81.1	385	25	US-09-649-163-1220	Sequence 1220, App	c 567	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 435	15.4	81.1	399	16	US-09-235-076-8801	Sequence 8801, App	c 568	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 436	15.4	81.1	399	16	US-09-248-797-27015	Sequence 27015, A	c 569	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 437	15.4	81.1	399	16	US-09-342-782-8801	Sequence 8801, App	c 570	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 438	15.4	81.1	399	29	US-09-737-223-8801	Sequence 8801, App	c 571	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 439	15.4	81.1	399	34	US-09-918-995-8801	Sequence 8801, App	c 572	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 440	15.4	81.1	401	25	US-09-925-544-27015	Sequence 27015, A	c 573	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 441	15.4	81.1	401	25	US-09-649-164-4251	Sequence 4251, App	c 574	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 442	15.4	81.1	407	16	US-09-269-618-43351	Sequence 43351, A	c 575	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 443	15.4	81.1	407	6	US-08-220-691-9331	Sequence 9331, App	c 576	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 444	15.4	81.1	409	6	US-08-220-691B-9331	Sequence 9331, App	c 577	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 445	15.4	81.1	409	6	US-08-220-691B-9331	Sequence 9331, App	c 578	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 446	15.4	81.1	422	24	US-09-620-111B-8304	Sequence 8304, App	c 579	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 447	15.4	81.1	430	22	US-09-553-093-6781	Sequence 6781, App	c 580	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 448	15.4	81.1	432	30	US-09-790-478-4754	Sequence 4754, App	c 581	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 449	15.4	81.1	436	25	US-09-649-163-6521	Sequence 6521, App	c 582	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 450	15.4	81.1	436	25	US-09-652-911-9190	Sequence 9190, App	c 583	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 451	15.4	81.1	436	40	US-09-770-171-2068	Sequence 2068, App	c 584	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 452	15.4	81.1	436	57	US-09-130-180-2107	Sequence 2107, App	c 585	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 453	15.4	81.1	438	61	US-60-177-571-1654	Sequence 1654, App	c 586	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 454	15.4	81.1	438	61	US-60-177-571-1654	Sequence 1654, App	c 587	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 455	15.4	81.1	443	61	US-60-177-571-1654	Sequence 1654, App	c 588	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 456	15.4	81.1	443	61	US-60-177-571-1654	Sequence 1654, App	c 589	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 457	15.4	81.1	447	25	US-09-654-617-237737	Sequence 237737, App	c 590	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 458	15.4	81.1	447	25	US-09-684-016-237737	Sequence 237737, App	c 591	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 459	15.4	81.1	447	32	US-09-708-427-42339	Sequence 42339, A	c 592	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 460	15.4	81.1	447	32	US-09-850-147-1373	Sequence 1373, A	c 593	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 461	15.4	81.1	447	64	US-60-202-213-1370	Sequence 1370, App	c 594	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 462	15.4	81.1	447	25	US-09-652-127-1381	Sequence 1381, App	c 595	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 463	15.4	81.1	449	25	US-09-654-617-454455	Sequence 454455, App	c 596	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 464	15.4	81.1	449	27	US-09-684-016-454455	Sequence 454455, App	c 597	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 465	15.4	81.1	497	6	US-08-220-661A-1077	Sequence 1077, App	c 598	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 466	15.4	81.1	497	6	US-08-220-661B-1077	Sequence 1077, App	c 599	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 467	15.4	81.1	497	6	US-08-220-662-1077	Sequence 1077, App	c 600	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 468	15.4	81.1	504	16	US-09-235-076-17326	Sequence 17326, A	c 601	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 469	15.4	81.1	504	16	US-09-237-227-15078	Sequence 15078, A	c 602	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 470	15.4	81.1	504	17	US-09-332-782-3725	Sequence 3725, A	c 603	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 471	15.4	81.1	504	29	US-09-737-223-37326	Sequence 37326, A	c 604	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 472	15.4	81.1	504	34	US-09-909-627-15078	Sequence 15078, A	c 605	15.4	81.1	528	14	US-09-668-632-1141	Sequence 1141, A
c 473	15.4	81.1	504	44	US-09-918-995-37326	Sequence 37326, A							

15.1	15.2	15.3	15.4	15.5	15.6	15.7	15.8	15.9	16.0	16.1	16.2	16.3	16.4	16.5	16.6	16.7	16.8	16.9	17.0	17.1	17.2	17.3	17.4	17.5	17.6	17.7	17.8	17.9	18.0	18.1	18.2	18.3	18.4	18.5	18.6	18.7	18.8	18.9	19.0	19.1	19.2	19.3	19.4	19.5	19.6	19.7	19.8	19.9	20.0	20.1	20.2	20.3	20.4	20.5	20.6	20.7	20.8	20.9	21.0	21.1	21.2	21.3	21.4	21.5	21.6	21.7	21.8	21.9	22.0	22.1	22.2	22.3	22.4	22.5	22.6	22.7	22.8	22.9	23.0	23.1	23.2	23.3	23.4	23.5	23.6	23.7	23.8	23.9	24.0	24.1	24.2	24.3	24.4	24.5	24.6	24.7	24.8	24.9	25.0	25.1	25.2	25.3	25.4	25.5	25.6	25.7	25.8	25.9	26.0	26.1	26.2	26.3	26.4	26.5	26.6	26.7	26.8	26.9	27.0	27.1	27.2	27.3	27.4	27.5	27.6	27.7	27.8	27.9	28.0	28.1	28.2	28.3	28.4	28.5	28.6	28.7	28.8	28.9	29.0	29.1	29.2	29.3	29.4	29.5	29.6	29.7	29.8	29.9	30.0	30.1	30.2	30.3	30.4	30.5	30.6	30.7	30.8	30.9	31.0	31.1	31.2	31.3	31.4	31.5	31.6	31.7	31.8	31.9	32.0	32.1	32.2	32.3	32.4	32.5	32.6	32.7	32.8	32.9	33.0	33.1	33.2	33.3	33.4	33.5	33.6	33.7	33.8	33.9	34.0	34.1	34.2	34.3	34.4	34.5	34.6	34.7	34.8	34.9	35.0	35.1	35.2	35.3	35.4	35.5	35.6	35.7	35.8	35.9	36.0	36.1	36.2	36.3	36.4	36.5	36.6	36.7	36.8	36.9	37.0	37.1	37.2	37.3	37.4	37.5	37.6	37.7	37.8	37.9	38.0	38.1	38.2	38.3	38.4	38.5	38.6	38.7	38.8	38.9	39.0	39.1	39.2	39.3	39.4	39.5	39.6	39.7	39.8	39.9	40.0	40.1	40.2	40.3	40.4	40.5	40.6	40.7	40.8	40.9	41.0	41.1	41.2	41.3	41.4	41.5	41.6	41.7	41.8	41.9	42.0	42.1	42.2	42.3	42.4	42.5	42.6	42.7	42.8	42.9	43.0	43.1	43.2	43.3	43.4	43.5	43.6	43.7	43.8	43.9	44.0	44.1	44.2	44.3	44.4	44.5	44.6	44.7	44.8	44.9	45.0	45.1	45.2	45.3	45.4	45.5	45.6	45.7	45.8	45.9	46.0	46.1	46.2	46.3	46.4	46.5	46.6	46.7	46.8	46.9	47.0	47.1	47.2	47.3	47.4	47.5	47.6	47.7	47.8	47.9	48.0	48.1	48.2	48.3	48.4	48.5	48.6	48.7	48.8	48.9	49.0	49.1	49.2	49.3	49.4	49.5	49.6	49.7	49.8	49.9	50.0	50.1	50.2	50.3	50.4	50.5	50.6	50.7	50.8	50.9	51.0	51.1	51.2	51.3	51.4	51.5	51.6	51.7	51.8	51.9	52.0	52.1	52.2	52.3	52.4	52.5	52.6	52.7	52.8	52.9	53.0	53.1	53.2	53.3	53.4	53.5	53.6	53.7	53.8	53.9	54.0	54.1	54.2	54.3	54.4	54.5	54.6	54.7	54.8	54.9	55.0	55.1	55.2	55.3	55.4	55.5	55.6	55.7	55.8	55.9	56.0	56.1	56.2	56.3	56.4	56.5	56.6	56.7	56.8	56.9	57.0	57.1	57.2	57.3	57.4	57.5	57.6	57.7	57.8	57.9	58.0	58.1	58.2	58.3	58.4	58.5	58.6	58.7	58.8	58.9	59.0	59.1	59.2	59.3	59.4	59.5	59.6	59.7	59.8	59.9	60.0	60.1	60.2	60.3	60.4	60.5	60.6	60.7	60.8	60.9	61.0	61.1	61.2	61.3	61.4	61.5	61.6	61.7	61.8	61.9	62.0	62.1	62.2	62.3	62.4	62.5	62.6	62.7	62.8	62.9	63.0	63.1	63.2	63.3	63.4	63.5	63.6	63.7	63.8	63.9	64.0	64.1	64.2	64.3	64.4	64.5	64.6	64.7	64.8	64.9	65.0	65.1	65.2	65.3	65.4	65.5	65.6	65.7	65.8	65.9	66.0	66.1	66.2	66.3	66.4	66.5	66.6	66.7	66.8	66.9	67.0	67.1	67.2	67.3	67.4	67.5	67.6	67.7	67.8	67.9	68.0	68.1	68.2	68.3	68.4	68.5	68.6	68.7	68.8	68.9	69.0	69.1	69.2	69.3	69.4	69.5	69.6	69.7	69.8	69.9	70.0	70.1	70.2	70.3	70.4	70.5	70.6	70.7	70.8	70.9	71.0	71.1	71.2	71.3	71.4	71.5	71.6	71.7	71.8	71.9	72.0	72.1	72.2	72.3	72.4	72.5	72.6	72.7	72.8	72.9	73.0	73.1	73.2	73.3	73.4	73.5	73.6	73.7	73.8	73.9	74.0	74.1	74.2	74.3	74.4	74.5	74.6	74.7	74.8	74.9	75.0	75.1	75.2	75.3	75.4	75.5	75.6	75.7	75.8	75.9	76.0	76.1	76.2	76.3	76.4	76.5	76.6	76.7	76.8	76.9	77.0	77.1	77.2	77.3	77.4	77.5	77.6	77.7	77.8	77.9	78.0	78.1	78.2	78.3	78.4	78.5	78.6	78.7	78.8	78.9	79.0	79.1	79.2	79.3	79.4	79.5	79.6	79.7	79.8	79.9	80.0	80.1	80.2	80.3	80.4	80.5	80.6	80.7	80.8	80.9	81.0	81.1	81.2	81.3	81.4	81.5	81.6	81.7	81.8	81.9	82.0	82.1	82.2	82.3	82.4	82.5	82.6	82.7	82.8	82.9	83.0	83.1	83.2	83.3	83.4	83.5	83.6	83.7	83.8	83.9	84.0	84.1	84.2	84.3	84.4	84.5	84.6	84.7	84.8	84.9	85.0	85.1	85.2	85.3	85.4	85.5	85.6	85.7	85.8	85.9	86.0	86.1	86.2	86.3	86.4	86.5	86.6	86.7	86.8	86.9	87.0	87.1	87.2	87.3	87.4	87.5	87.6	87.7	87.8	87.9	88.0	88.1	88.2	88.3	88.4	88.5	88.6	88.7	88.8	88.9	89.0	89.1	89.2	89.3	89.4	89.5	89.6	89.7	89.8	89.9	90.0	90.1	90.2	90.3	90.4	90.5	90.6	90.7	90.8	90.9	91.0	91.1	91.2	91.3	91.4	91.5	91.6	91.7	91.8	91.9	92.0	92.1	92.2	92.3	92.4	92.5	92.6	92.7	92.8	92.9	93.0	93.1	93.2	93.3	93.4	93.5	93.6	93.7	93.8	93.9	94.0	94.1	94.2	94.3	94.4	94.5	94.6	94.7	94.8	94.9	95.0	95.1	95.2	95.3	95.4	95.5	95.6	95.7	95.8	95.9	96.0	96.1	96.2	96.3	96.4	96.5	96.6	96.7	96.8	96.9	97.0	97.1	97.2	97.3	97.4	97.5	97.6	97.7	97.8	97.9	98.0	98.1	98.2	98.3	98.4	98.5	98.6	98.7	98.8	98.9	99.0	99.1	99.2	99.3	99.4	99.5	99.6	99.7	99.8	99.9	100.0
------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	------	-------

[illegible]





```

1 GENERAL INFORMATION:
2 Sequence: 30551, Application US/09542415
3 GENERAL INFORMATION:
4 APPLICANT: Sellhamer, Jeffrey J.
5 APPLICANT: Delecano, Angelo M.
6 APPLICANT: Stuart, Susan G.
7 APPLICANT: Stuve, Laura L.
8 APPLICANT: Mullaby, Sara J.
9 APPLICANT: Naughton, Rebecca E.
10 TITLE OF INVENTION: POLYMERIZABLES
11 FILE REFERENCE: PD-1002 CIP
12 CURRENT APPLICATION NUMBER: US/09542415
13 CURRENT FILING DATE: 2000-04-24
14 Prior application data removed - refer to PALM or file wrapper
15 NUMBER OF SEQ ID NOS: 14562
16 SOFTWARE: PERL Program
17 SEQ ID NO: 14044
18 LENGTH: 264
19 TYPE: DNA
20 ORGANISM: Homo sapiens
21 FEATURE:
22 NAME/KEY: misc-feature
23 OTHER INFORMATION: Incyte ID No: h001296954
24 US-09-540-229-14044
25
26 Query Match: 100.0%; Score 19; DB Z1: Length 264;
27 Best Local Similarity: 100.0%; Prod. No. 1,540-02;
28 Matches 19; Conservative 0; Mismatches 0; Indels 0
29
30 Q1 1 GGAAGAGGAGTGAAGG 19
31 19 GGAAGAGGAGTGAAGG 19
32
33 RESULT 5
34 US-09-010-256-11717
35 Sequence 1171, Application US/09-010-256
36 GENERAL INFORMATION:
37 APPLICANT: Goodrich, Joseph H.
38 APPLICANT: Stuart, Susan G.
39 APPLICANT: Stuve, Laura L.
40 APPLICANT: Mullaby, Sara J.
41 APPLICANT: Naughton, Rebecca E.
42 APPLICANT: Kline, David M.
43 TITLE OF INVENTION: POLYMERIZABLES
44 FILE REFERENCE: PD-1002 CIP
45 CURRENT APPLICATION NUMBER: US/09542415
46 CURRENT FILING DATE: 2000-04-24
47 Prior application data removed - refer to PALM or file wrapper
48 NUMBER OF SEQ ID NOS: 42212
49 SOFTWARE: PERL Program
50 SEQ ID NO: 40444
51 LENGTH: 256
52 TYPE: DNA
53 ORGANISM: Homo sapiens
54 FEATURE:
55 NAME/KEY: misc-feature
56 OTHER INFORMATION: Incyte ID No: h001166289
57 US-09-532-415-40444
58
59 Query Match: 100.0%; Score 19; DB Z1: Length 256;
60 Best Local Similarity: 100.0%; Prod. No. 1,540-02;
61 Matches 19; Conservative 0; Mismatches 0; Indels 0
62
63 Q1 1 GGAAGAGGAGTGAAGG 19
64 19 GGAAGAGGAGTGAAGG 19
65
66 RESULT 5
67 US-09-532-415-40444
68 Sequence 044, Application US/09542415
69 GENERAL INFORMATION:
70 APPLICANT: Sellhamer, Jeffrey J.
71 APPLICANT: Delecano, Angelo M.
72 APPLICANT: Stuart, Susan G.
73 APPLICANT: Stuve, Laura L.
74 APPLICANT: Mullaby, Sara J.
75 APPLICANT: Naughton, Rebecca E.
76 TITLE OF INVENTION: POLYMERIZABLES
77 FILE REFERENCE: PD-1002 CIP
78 CURRENT APPLICATION NUMBER: US/09542415
79 CURRENT FILING DATE: 2000-04-24
80 Prior application data removed - refer to PALM or file wrapper
81 NUMBER OF SEQ ID NOS: 42212
82 SOFTWARE: PERL Program
83 SEQ ID NO: 40444
84 LENGTH: 256
85 TYPE: DNA
86 ORGANISM: Homo sapiens
87 FEATURE:
88 NAME/KEY: misc-feature
89 OTHER INFORMATION: Incyte ID No: h001166289
90 US-09-532-415-40444
91
92 Query Match: 100.0%; Score 19; DB Z1: Length 256;
93 Best Local Similarity: 100.0%; Prod. No. 1,540-02;
94 Matches 19; Conservative 0; Mismatches 0; Indels 0
95
96 Q1 1 GGAAGAGGAGTGAAGG 19
97 19 GGAAGAGGAGTGAAGG 19
98
99 RESULT 4
100 US-09-540-229-14044
101 Sequence 14044, Application US/09540229

```

```

1 GENERAL INFORMATION:
2 Sequence: 30551, Application US/09542415
3 GENERAL INFORMATION:
4 APPLICANT: Sellhamer, Jeffrey J.
5 APPLICANT: Delecano, Angelo M.
6 APPLICANT: Stuart, Susan G.
7 APPLICANT: Stuve, Laura L.
8 APPLICANT: Mullaby, Sara J.
9 APPLICANT: Naughton, Rebecca E.
10 TITLE OF INVENTION: POLYMERIZABLES
11 FILE REFERENCE: PD-1002 CIP
12 CURRENT APPLICATION NUMBER: US/09542415
13 CURRENT FILING DATE: 2000-04-24
14 Prior application data removed - refer to PALM or file wrapper
15 NUMBER OF SEQ ID NOS: 42212
16 SOFTWARE: PERL Program
17 SEQ ID NO: 40451
18 LENGTH: 198
19 TYPE: DNA
20 ORGANISM: Homo sapiens
21 FEATURE:
22 NAME/KEY: misc-feature
23 OTHER INFORMATION: Incyte ID No: h001296954
24 US-09-532-415-40351
25
26 Query Match: 100.0%; Score 19; DB Z1: Length 198;
27 Best Local Similarity: 100.0%; Prod. No. 1,540-02;
28 Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
29
30 Q1 1 GGAAGAGGAGTGAAGG 19
31 19 GGAAGAGGAGTGAAGG 31
32
33 RESULT 5
34 US-09-532-415-40444
35 Sequence 044, Application US/09542415
36 GENERAL INFORMATION:
37 APPLICANT: Sellhamer, Jeffrey J.
38 APPLICANT: Delecano, Angelo M.
39 APPLICANT: Stuart, Susan G.
40 APPLICANT: Stuve, Laura L.
41 APPLICANT: Mullaby, Sara J.
42 APPLICANT: Naughton, Rebecca E.
43 TITLE OF INVENTION: POLYMERIZABLES
44 FILE REFERENCE: PD-1002 CIP
45 CURRENT APPLICATION NUMBER: US/09542415
46 CURRENT FILING DATE: 2000-04-24
47 Prior application data removed - refer to PALM or file wrapper
48 NUMBER OF SEQ ID NOS: 42212
49 SOFTWARE: PERL Program
50 SEQ ID NO: 40444
51 LENGTH: 256
52 TYPE: DNA
53 ORGANISM: Homo sapiens
54 FEATURE:
55 NAME/KEY: misc-feature
56 OTHER INFORMATION: Incyte ID No: h001166289
57 US-09-532-415-40444
58
59 Query Match: 100.0%; Score 19; DB Z1: Length 256;
60 Best Local Similarity: 100.0%; Prod. No. 1,540-02;
61 Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
62
63 Q1 1 GGAAGAGGAGTGAAGG 19
64 19 GGAAGAGGAGTGAAGG 77
65
66 RESULT 4
67 US-09-540-229-14044
68 Sequence 14044, Application US/09540229

```



us-09-925-548-6.rnrm

Sun May 25 14:51:39 2003

```

1  NUMBER OF SEQUENCES: 1746
2  C-RESPONSE: ADDRESS: 1746
3  ADDRESSEE: 16-YE FARMER, BENTLEY, 1746
4  STREET: 4174 PORTER DRIVE
5  CITY: PALO ALTO
6  STATE: CALIFORNIA
7  COUNTRY: USA
8  ZIP: 94304
9  NUMBER RELEASABLE FORM:
10  FILING DATE: 1999-01-24
11  FILING TYPE: Floppy disk
12  OPERATING SYSTEM: IBM PC compatible
13  SOFTWARE: WORD Perfect 6.1.1 for Windows/MS-DOS 4.2
14  CURRENT APPLICATION DATA:
15  CLASSIFICATION:
16  NAME: GERRONE, MICHAEL C.
17  REGISTRATION NUMBER: 39,142
18  REFERENCE NUMBER: PD-0406P
19  TELECOMMUNICATION INFORMATION:
20  TELEPHONE: (415) 855-0555
21  TELEFAX: (415) 845-4166
22  INFORMATION FOR SEQ ID NO: 1960:
23  SEQUENCE CHARACTERISTICS:
24  LENGTH: 404 base pairs
25  TYPE: nucleic acid
26  STRANDEDNESS: single
27  TOPOLOGY: linear
28  Molecule type: cDNA
29  IMMEDIATE SOURCE:
30  CLONE: 253045H1
31  US-09-085-485-1060

```

```

Query Match 100.0% Score 19; DB 47; Length 404;
Best Local Similarity 100.0%; Pred. No. 1,600-02;
Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 1 GGGAGAGAGATGAAGCC 19
DB 29 GGGAGAGAGATGAAGCC 11

```

```

1  US-09-24-611-12842/c
2  Sequence 12842, Application US/09244611
3  GENERAL INFORMATION:
4  APPLICANT: Hyseq, Inc.
5  TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
6  FILE REFERENCE: 20411-757
7  CURRENT FILING DATE: 1999-01-21
8  NUMBER OF SEQ ID NOS: 21025
9  SOFTWARE: FastSeq for Windows Version 4.0
10  SEQ ID NO: 12842
11  LENGTH: 424
12  TYPE: DNA
13  ORGANISM: Homo sapiens
14  US-09-24-611-12842

```

```

Query Match 100.0% Score 19; DB 16; Length 424;
Best Local Similarity 100.0%; Pred. No. 1,460-02;
Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 1 GGGAGAGAGATGAAGCC 19
DB 45 GGGAGAGAGATGAAGCC 17

```

```

1  US-09-284-768-17564/c
2  Sequence 17564, Application US/09284768
3  GENERAL INFORMATION:
4  APPLICANT: Hyseq, Inc.
5  TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
6  FILE REFERENCE: 20411-757
7  CURRENT FILING DATE: 1999-01-21
8  NUMBER OF SEQ ID NOS: 21025
9  SOFTWARE: FastSeq for Windows Version 4.0
10  SEQ ID NO: 17564
11  LENGTH: 424
12  TYPE: DNA
13  ORGANISM: Homo sapiens
14  US-09-284-768-17564

```

```

1  Sequence 17564, Application US/09284768
2  GENERAL INFORMATION:
3  APPLICANT: Hyseq, Inc.
4  TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
5  FILE REFERENCE: 20411-757
6  CURRENT FILING DATE: 1999-01-21
7  NUMBER OF SEQ ID NOS: 21025
8  SOFTWARE: FastSeq for Windows Version 4.0
9  SEQ ID NO: 17564
10  LENGTH: 424
11  TYPE: DNA
12  ORGANISM: Homo sapiens
13  US-09-284-768-17564

```

```

Query Match 100.0% Score 19; DB 16; Length 424;
Best Local Similarity 100.0%; Pred. No. 1,460-02;
Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 1 GGGAGAGAGATGAAGCC 19
DB 45 GGGAGAGAGATGAAGCC 17

```

```

1  US-09-004-809-12842/c
2  Sequence 12842, Application US/09004809
3  GENERAL INFORMATION:
4  APPLICANT: Hyseq, Inc.
5  TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
6  FILE REFERENCE: 20411-757
7  CURRENT FILING DATE: 2001-07-12
8  NUMBER OF SEQ ID NOS: 21025
9  SOFTWARE: FastSeq for Windows Version 4.0
10  SEQ ID NO: 12842
11  LENGTH: 424
12  TYPE: DNA
13  ORGANISM: Homo sapiens
14  US-09-004-809-12842

```

```

Query Match 100.0% Score 19; DB 47; Length 424;
Best Local Similarity 100.0%; Pred. No. 1,600-02;
Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 1 GGGAGAGAGATGAAGCC 19
DB 45 GGGAGAGAGATGAAGCC 17

```

```

1  US-09-909-897-17564/c
2  Sequence 17564, Application US/0909897
3  GENERAL INFORMATION:
4  APPLICANT: Hyseq, Inc.
5  TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
6  FILE REFERENCE: 20411-757
7  CURRENT FILING DATE: 2001-08-21
8  NUMBER OF SEQ ID NOS: 21025
9  SOFTWARE: FastSeq for Windows Version 4.0
10  SEQ ID NO: 17564
11  LENGTH: 424
12  TYPE: DNA
13  ORGANISM: Homo sapiens
14  US-09-909-897-17564

```

```

Query Match 100.0% Score 19; DB 16; Length 424;
Best Local Similarity 100.0%; Pred. No. 1,460-02;
Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 1 GGGAGAGAGATGAAGCC 19
DB 45 GGGAGAGAGATGAAGCC 17

```

```

1  US-09-909-897-17564/c
2  Sequence 17564, Application US/0909897
3  GENERAL INFORMATION:
4  APPLICANT: Hyseq, Inc.
5  TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
6  FILE REFERENCE: 20411-757
7  CURRENT FILING DATE: 2001-08-21
8  NUMBER OF SEQ ID NOS: 21025
9  SOFTWARE: FastSeq for Windows Version 4.0
10  SEQ ID NO: 17564
11  LENGTH: 424
12  TYPE: DNA
13  ORGANISM: Homo sapiens
14  US-09-909-897-17564

```







```

? CURRENT FILING DATE: 2000-08-30
? PRIOR APPLICATION NUMBER: 62/151,136
? PRIOR FILING DATE: 1999-08-30
? NUMBER OF SEQ ID NOS: 11227
? SOFTWARE: FASTSEQ for Windows Version 4.0
? SEQ ID NO 4014
? LENGTH: 441
? TYPE: DNA
? ORGANISM: Homo sapiens
? FEATURE:
? NAME/KEY: misc_feature
? LOCATION: (1) ... (441)
? OTHER INFORMATION: n A, T, C or G
US-09-925-548-6.rnmpm

Query Match
Best Local Similarity 100.0%; Score 19; DB 25; Length 441;
Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GGGAAAGACGATGAAACCC 19
DB 124 GGGAAAGACGATGAAACCC 106

RESULT 26
US-09-925-548-6.rnmpm
? Sequence 2179, Application US/09/09720
? GENERAL INFORMATION:
? APPLICANT: Rescal, Inc.
? TITLE OF INVENTION: Novel Nucleic Acid Sequences Obtained
? FILE REFERENCE: 20411-777
? CURRENT APPLICATION NUMBER: 92/599,720
? CURRENT FILING DATE: 1999-09-21
? NUMBER OF SEQ ID NOS: 20869
? SOFTWARE: Hy-patent.pl Version 3.1
? SEQ ID NO 2179
? LENGTH: 449
? TYPE: DNA
? ORGANISM: Homo sapiens
? FEATURE:
? NAME/KEY: misc_feature
? LOCATION: (1) ... (449)
? OTHER INFORMATION: n A, T, C or G
US-09-925-548-6.rnmpm

Query Match
Best Local Similarity 100.0%; Score 19; DB 17; Length 449;
Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GGGAAAGACGATGAAACCC 19
DB 389 GGGAAAGACGATGAAACCC 421

RESULT 27
US-09-925-548-6.rnmpm
? Sequence 2179, Application US/09/21478
? GENERAL INFORMATION:
? APPLICANT: Hyseq, Inc.
? TITLE OF INVENTION: Novel Nucleic Acid Sequences Obtained
? FILE REFERENCE: 20411-777
? CURRENT APPLICATION NUMBER: 92/599,721, 378
? CURRENT FILING DATE: 2001-08-02
? PRIOR FILING DATE: 1999-09-21
? NUMBER OF SEQ ID NOS: 20869
? SOFTWARE: Hy-patent.pl Version 4.1
? SEQ ID NO 2179
? LENGTH: 449
? TYPE: DNA

```

```

? ORGANISM: Homo sapiens
? FEATURE:
? NAME/KEY: misc_feature
? LOCATION: (1) ... (449)
? OTHER INFORMATION: n A, T, C or G
US-09-925-548-6.rnmpm

Query Match
Best Local Similarity 100.0%; Score 19; DB 54; Length 449;
Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GGGAAAGACGATGAAACCC 19
DB 389 GGGAAAGACGATGAAACCC 421

RESULT 28
US-09-925-548-6.rnmpm
? Sequence 2179, Application US/09/09720
? GENERAL INFORMATION:
? APPLICANT: Rescal, Inc.
? TITLE OF INVENTION: Novel Nucleic Acid Sequences Obtained
? FILE REFERENCE: 20411-777
? CURRENT APPLICATION NUMBER: 92/599,720
? CURRENT FILING DATE: 1999-09-21
? EARLIER FILING DATE: 1998-09-14
? EARLIER FILING DATE: 1998-09-14
? EARLIER FILING DATE: 1998-11-05
? EARLIER FILING DATE: 1998-11-05
? EARLIER FILING DATE: 1999-04-30
? NUMBER OF SEQ ID NOS: 5229
? SOFTWARE: Hy-patent.pl Version 3.1
? SEQ ID NO 2179
? LENGTH: 449
? TYPE: DNA
? ORGANISM: Homo sapiens
US-09-925-548-6.rnmpm

Query Match
Best Local Similarity 100.0%; Score 19; DB 17; Length 449;
Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GGGAAAGACGATGAAACCC 19
DB 389 GGGAAAGACGATGAAACCC 421

RESULT 29
US-09-925-548-6.rnmpm
? Sequence 2179, Application US/09/21478
? GENERAL INFORMATION:
? APPLICANT: Rescal, Inc.
? APPLICANT: Rescal, Inc.
? APPLICANT: Rescal, Inc.
? TITLE OF INVENTION: Novel Nucleic Acid Sequences Obtained
? FILE REFERENCE: 20411-777
? CURRENT APPLICATION NUMBER: 92/599,720
? CURRENT FILING DATE: 2001-08-02
? PRIOR FILING DATE: 1999-09-21
? NUMBER OF SEQ ID NOS: 20869
? SOFTWARE: Hy-patent.pl Version 4.1
? SEQ ID NO 2179
? LENGTH: 449
? TYPE: DNA
? ORGANISM: Homo sapiens
? FEATURE:
? NAME/KEY: misc_feature
? LOCATION: (1) ... (449)

```









```

: TYPE: DNA
: ORGANISM: Homo sapiens
: FEATURE:
: NAME/KEY: misc_feature
: OTHER INFORMATION: Intron ID No: 474175.41
US-09-925-548-600-10610

```

```

Query Match 100.0% Score 19: 106 67: Length 764
Best Local Similarity 100.0% Pred. No: 1,80-02
Matches 19: Conservative 0: Mismatches 0: Indels 0: Gaps 0:

```

```

QY 1 GGAAGAGAGATGAACGCC 19
ID 479 GGAAGAGAGATGAACGCC 461

```

```

RESULT 42
US-09-925-548-600-10610
: Sequence 600, Applied to us/09-925-548-600-10610
: GENERAL INFORMATION:
: APPLICANT: Morris, MacDonald
: TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: FILE REFERENCE: CX-0021-1 P
: CURRENT FILING DATE: 2000-06-21
: PRIOR APPLICATION NUMBER: 60-2229
: NUMBER OF SEQ ID NOS: 847
: SEQ ID NO 10610
: TYPE: DNA
: LENGTH: 764
: ORGANISM: Homo sapiens
: FEATURE:
: NAME/KEY: misc_feature
: OTHER INFORMATION: Intron ID No: 474175.41
US-09-925-548-600-10610

```

```

Query Match 100.0% Score 19: 106 67: Length 764
Best Local Similarity 100.0% Pred. No: 1,80-02
Matches 19: Conservative 0: Mismatches 0: Indels 0: Gaps 0:

```

```

QY 1 GGAAGAGAGATGAACGCC 19
ID 479 GGAAGAGAGATGAACGCC 461

```

```

RESULT 43
US-09-925-548-600-2229
: Sequence 2229, Applied to us/09-925-548-600-2229
: GENERAL INFORMATION:
: APPLICANT: Morris, MacDonald
: TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: FILE REFERENCE: CX-0014 P
: CURRENT FILING DATE: 2000-06-21
: PRIOR APPLICATION NUMBER: 60-2229
: NUMBER OF SEQ ID NOS: 847
: SEQ ID NO 2229
: TYPE: DNA
: LENGTH: 764
: ORGANISM: Homo sapiens
: FEATURE:
: NAME/KEY: misc_feature
: OTHER INFORMATION: Intron ID No: 995745.1
US-09-925-548-600-2229

```

```

Query Match 100.0% Score 19: 106 67: Length 764
Best Local Similarity 100.0% Pred. No: 1,80-02
Matches 19: Conservative 0: Mismatches 0: Indels 0: Gaps 0:

```

```

QY 1 GGAAGAGAGATGAACGCC 19
ID 479 GGAAGAGAGATGAACGCC 461

```

```

RESULT 44
US-09-925-548-600-2229
: Sequence 2229, Applied to us/09-925-548-600-2229
: GENERAL INFORMATION:
: APPLICANT: Morris, MacDonald
: TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: FILE REFERENCE: CX-0014 P
: CURRENT FILING DATE: 2000-06-21
: PRIOR APPLICATION NUMBER: 60-2229
: NUMBER OF SEQ ID NOS: 847
: SEQ ID NO 2229
: TYPE: DNA
: LENGTH: 764
: ORGANISM: Homo sapiens
: FEATURE:
: NAME/KEY: misc_feature
: OTHER INFORMATION: Intron ID No: 995745.1
US-09-925-548-600-2229

```

```

Query Match 100.0% Score 19: 106 67: Length 764
Best Local Similarity 100.0% Pred. No: 1,80-02
Matches 19: Conservative 0: Mismatches 0: Indels 0: Gaps 0:

```

```

QY 1 GGAAGAGAGATGAACGCC 19
ID 479 GGAAGAGAGATGAACGCC 461

```

```

RESULT 44
US-09-925-548-600-2229
: Sequence 2229, Applied to us/09-925-548-600-2229
: GENERAL INFORMATION:
: APPLICANT: Morris, MacDonald
: TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: FILE REFERENCE: CX-0014 P
: CURRENT FILING DATE: 2000-06-21
: PRIOR APPLICATION NUMBER: 60-2229
: NUMBER OF SEQ ID NOS: 847
: SEQ ID NO 2229
: TYPE: DNA
: LENGTH: 764
: ORGANISM: Homo sapiens
: FEATURE:
: NAME/KEY: misc_feature
: OTHER INFORMATION: Intron ID No: 995745.1
US-09-925-548-600-2229

```

```

Query Match 100.0% Score 19: 106 67: Length 764
Best Local Similarity 100.0% Pred. No: 1,80-02
Matches 19: Conservative 0: Mismatches 0: Indels 0: Gaps 0:

```

```

QY 1 GGAAGAGAGATGAACGCC 19
ID 479 GGAAGAGAGATGAACGCC 461

```

```

RESULT 45
US-09-925-548-600-2229
: Sequence 2229, Applied to us/09-925-548-600-2229
: GENERAL INFORMATION:
: APPLICANT: Morris, MacDonald
: TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: FILE REFERENCE: CX-0014 P
: CURRENT FILING DATE: 2000-06-21
: PRIOR APPLICATION NUMBER: 60-2229
: NUMBER OF SEQ ID NOS: 847
: SEQ ID NO 2229
: TYPE: DNA
: LENGTH: 764
: ORGANISM: Homo sapiens
: FEATURE:
: NAME/KEY: misc_feature
: OTHER INFORMATION: Intron ID No: 995745.1
US-09-925-548-600-2229

```

```

Query Match 100.0% Score 19: 106 67: Length 764
Best Local Similarity 100.0% Pred. No: 1,80-02
Matches 19: Conservative 0: Mismatches 0: Indels 0: Gaps 0:

```

```

QY 1 GGAAGAGAGATGAACGCC 19
ID 479 GGAAGAGAGATGAACGCC 461

```

```

RESULT 46
US-09-925-548-600-2229
: Sequence 2229, Applied to us/09-925-548-600-2229
: GENERAL INFORMATION:
: APPLICANT: Morris, MacDonald
: TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: TITLE OF INVENTOR: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE POLYMER
: FILE REFERENCE: CX-0014 P
: CURRENT FILING DATE: 2000-06-21
: PRIOR APPLICATION NUMBER: 60-2229
: NUMBER OF SEQ ID NOS: 847
: SEQ ID NO 2229
: TYPE: DNA
: LENGTH: 764
: ORGANISM: Homo sapiens
: FEATURE:
: NAME/KEY: misc_feature
: OTHER INFORMATION: Intron ID No: 995745.1
US-09-925-548-600-2229

```



Sun May 25 14:51:39 2003

```

: TITLE: FILING DATE: 1999-03-19
:
: NUMBER: F SEQ ID NOS: 202
:
: SLEEPWAKE: Part out in Ver. 2.1
: SEQ ID NO: 1-3
:
: LENGTH: 1780
:
: TYPE: tNA
:
: ORGANISM: Homo sapiens
:
: 1-3, 1-3, 4-10-13

```

Query Match	100.00;	Score 19;	100.40;	Length 1780;
Best local Similarity	100.00;	Pred. No. 20.02;		
Matches 19;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;

[illegible]

Search completed; May 23, 2003, 02:59:44  
Job time: 2123 secs

